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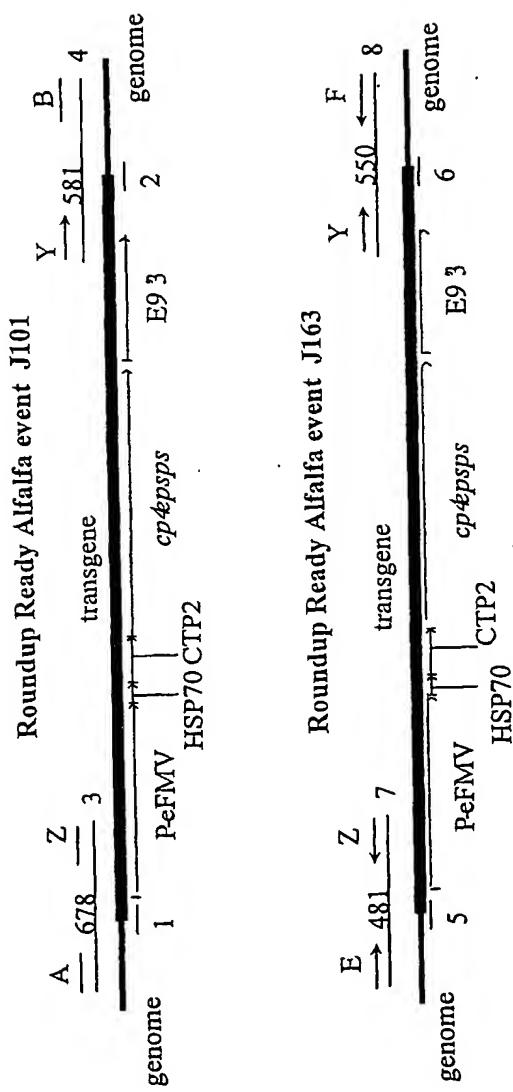


FIGURE 1

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1 ACATAGCTTT TATGTGAAGG AAAAATCAAA TTTTCCAAAAA TTTGGAGTTT  
51 ATAGACTGAG CACATGATAC TGTCGGTGTT TGTTTAAAG ATTAAAAAAC  
101 TATAACCCTTT GAATAATTAA ATTAAATCAA ATTTTCATAT TAAATTTTA  
151 ATTTTATAGT AATAATCTAA TTTTAATACA CTTAGGTGTA AAAAAAAATTT  
201 AAGCTTCAAA GTTTATATT GTCAGCAAAT CACAACTAAT TGTGTGTACG  
251 GATACAAAGT CAAACATGAT TTATTGACGG TGTAAAAAAAT CTTTACAGTG  
301 ACAATGTATA TGGATTAAAT CGATTTATA TTAGTTATTT TATGTTATAT  
351 CGTATTTCATG TCATGTGTTT TGTACTGATC TTGTGTCTATA GTTCAAAACA  
401 CTGATAGTTT AACTGAAGG CGGGAAACGA CAATCTGATC CCCATCAAGC  
451 TTCTGCAGGT CCTGCTCGAG TGGAAGCTAA TTCTCAGTCC AAAGCCTCAA  
501 CAAGGTCAGG GTACAGAGTC TCCAAACCAT TAGCCAAAAG CTACAGGAGA  
551 TCAATGAAGA ATCTTCATC AAAGTAAACT ACTGTTCCAG CACATGCATC  
601 ATGGTCAGTA AGTTTCAGAA AAAGACATCC ACCGAAGACT TAAAGTTAGT  
651 GGGCATCTTT GAAAGTAATC TTGTCAAC

FIGURE 2

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1 GAACTTTCCT TTATGTAATTTCCAGAATC CTTGTCAGAT TCTAATCATT  
51 GCTTATAAT TATAGTATA CTCATGGATT TGTAGTTGAG TATGAAAATA  
101 TTTTTAATG CATTATGA CTTGCCAATT GATTGACAAC ATGCATCAAT  
151 CGACCTGCAG CCAC~~T~~CGAAG CGGCCGCCAC TCGAGTGGTG GCCGCATCGA  
201 TCGTGAAGTT TCTCATCTAA GCCCCATTT GGACGTGAAT GTAGACACGT  
251 CGAAATAAAG ATTTCCGAAT TAGAATAATT TGTTTATTGC TTTCGCCTAT  
301 AAATACGACG GATCGTATGC ATTAATATA TAGAGGAATT TCTTATCTTG  
351 CTAATTCCAG CATAGTTATT TTAATTGTC AAATAATTGT ATGAATGGTA  
401 TATTCTACAA CTTTCTCATG CTTTCAATA ATGATGTTGT TTTCTGTAA  
451 TGCTATTTG ATTTTATTT GCAGGTGAAC AAGAGCAACA GTGCTTGAT  
501 CCATATTACA AATGAGGGAT GGTACTATT TATGGATGCA GTGAAACTTC  
551 TGAAAACATA GTTGGTGTGG TGCTGTTGGA G

FIGURE 3

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1 TTTCTTACTT CCTAGATTTG GTAGGGTTGA AACACATGAA AATTTAAAGC  
51 ATATACACAA TACATTTGG ACTTGACTTG ACTTCACTAT TCATAATGAG  
101 CTTCATGCAT ATTTGGATGC CCATATCATA TCAAGTCATT ATTTTATTT  
151 CCTTTAACG ATTACCCCT CCTACTTTT TCCTTCTTG CCGGGACAAG  
201 GTCATCCAAA CTGAAGTGTG CGGTGGAAA CGACACTCTG ATCCCCATCA  
251 AGCTTCTGCA GGTCTGCTC GAGTGGAAAGC TAATTCTAG TCCAAAGCCT  
301 CAACAAGGTC AGGGTACAGA GTCTCCAAAC CATTAGCCAA AAGCTACAGG  
351 AGATCAATGA AGAATCTTCA ATCAAAGTAA ACTACTGTTC CAGCACATGC  
401 ATCATGGTCA GTAAGTTCA GAAAAAGACA TCCACCGAAG ACTTAAAGTT  
451 AGTGGGCATC TTTGAAAGTA ATCTTGTCAA C

FIGURE 4

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1 GAACTTCCT TTATGTAATT TTCCAGAATC CTTGTCAGAT TCTAATCATT  
51 GCTTTATAAT TATAGTATA CTCATGGATT TGTAGTTGAG TATGAAAATA  
101 TTTTTAATG CATTATGA CTTGCCAATT GATTGACAAC ATGCATCAAT  
151 CGACCTGAG CGACTCGAAG CGGCCGCCAC TCGAGTGGTG GCCGCATCGA  
201 TCGTGAAGTT TCTCATCTAA GCCCCATTG GGACGTGAAT GTAGACACGT  
251 CGAAATAAAG ATTTCCAAT TAGAATAATT TGTTTATTGC TTTGCCATAT  
301 AAATACGACG GATCGTAATT TGTCGTTTA TCAAAATGTA CTTTCATT  
351 ATAATAACTT CCATTTTTT TTTCTTTTC TTTTATAATA ACAGAAAAAG  
401 AAAAAGAAAG ATGATGAAAA GAGAAAAGAG AAAACCGAAC CATGATAATT  
451 AACACACCAC GTGCAATTAA CTTTACTTTA ATTTACTAC TACCTTATT  
501 TTTCTTCAGC GTGGTAACCG TTATACTCTT TATTACACCA CTCACCCACCA

FIGURE 5

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A 5' ACATAGCTTTATGTGAAGGAAAAATC 3' (SEQ ID NO:9)  
E 5' TTTCTTACTTCCTAGATTTGGTAGG 3' (SEQ ID NO:10)  
Z 5' GTTGACAAGATTACTTCAAAGATGC 3' (SEQ ID NO:11)  
B 5' CTCCAACAGCACCCACACCAACTAT 3' (SEQ ID NO:12)  
F 5' TGGTGGTGAGTGGTGTAAATAAGAG 3' (SEQ ID NO:13)  
Y 5' GAACTTTCCCTTATGTAATTTCAG 3' (SEQ ID NO:14)

FIGURE 6

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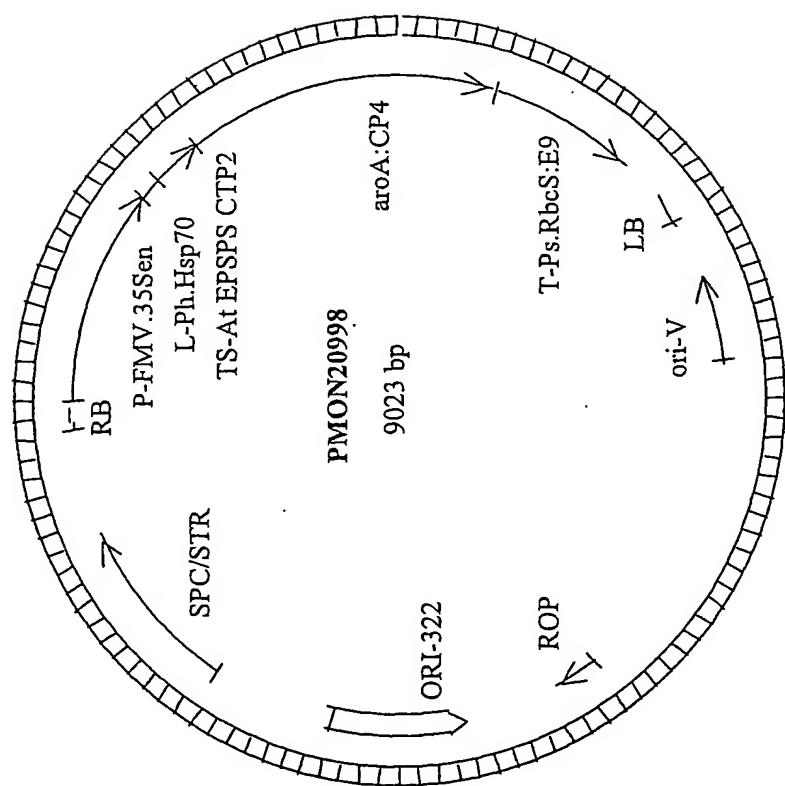


FIGURE 7